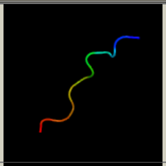
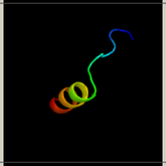
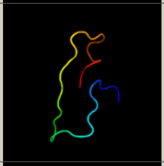
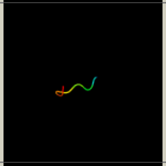
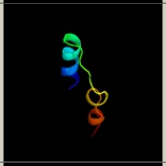
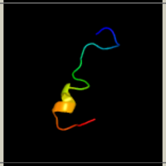
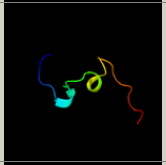
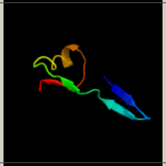
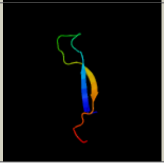
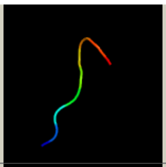
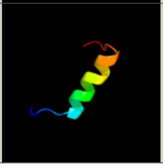
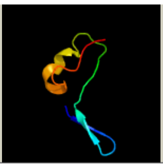
Email	mdejesus@rockefeller.edu
Description	RVBD2348c_(-)_2626664_2626990
Date	Mon Aug 5 13:25:50 BST 2019
Unique Job ID	dfbe29c7a0d1582f

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c6hm3B_	Alignment		28.3	67	**PDB header:**cell cycle **Chain:** B: **PDB Molecule:**dna replication regulator sld3; **PDBTitle:** crystal structure of rad4 brct1,2 in complex with a sld32 phosphopeptide
2	c2m4hA_	Alignment		20.4	60	**PDB header:**viral protein **Chain:** A: **PDB Molecule:**feline calicivirus vpg protein; **PDBTitle:** solution structure of the core domain (10-76) of the feline2 calicivirus vpg protein
3	c2mxdA_	Alignment		16.2	53	**PDB header:**viral protein **Chain:** A: **PDB Molecule:**viral protein genome-linked; **PDBTitle:** solution structure of vpg of porcine sapovirus
4	c3acgA_	Alignment		13.1	45	**PDB header:**hydrolase **Chain:** A: **PDB Molecule:**beta-1,4-endoglucanase; **PDBTitle:** crystal structure of carbohydrate-binding module family 28 from2 clostridium josui cel5a in complex with cellobiose
5	d1uwwa_	Alignment		12.3	45	**Fold:**Galactose-binding domain-like **Superfamily:**Galactose-binding domain-like **Family:**Family 28 carbohydrate binding module, CBM28
6	c2xumS_	Alignment		10.0	78	**PDB header:**oxidoreductase/peptide **Chain:** S: **PDB Molecule:**asp-substrate peptide 2; **PDBTitle:** factor inhibiting hif (fih) q239h mutant in complex with zn(ii), nog2 and asp-substrate peptide (20-mer)
7	c3bv8A_	Alignment		9.8	24	**PDB header:**transferase **Chain:** A: **PDB Molecule:**tetrahydrodipicolinate acetyltransferase; **PDBTitle:** crystal structure of the n-terminal domain of tetrahydrodipicolinate2 acetyltransferase from staphylococcus aureus
8	c2rqyA_	Alignment		9.6	24	**PDB header:**unknown function **Chain:** A: **PDB Molecule:**putative uncharacterized protein; **PDBTitle:** solution structure and dynamics of mouse armet
9	d2buda1	Alignment		8.3	21	**Fold:**SH3-like barrel **Superfamily:**Chromo domain-like **Family:**Chromo barrel domain
10	d2b3ya2	Alignment		8.1	32	**Fold:**Aconitase iron-sulfur domain **Superfamily:**Aconitase iron-sulfur domain **Family:**Aconitase iron-sulfur domain
11	d1ffkf_	Alignment		7.5	13	**Fold:**alpha/beta-Hammerhead **Superfamily:**Ribosomal protein L16p/L10e **Family:**Ribosomal protein L10e

12	c5w54A_	Alignment		7.2	67	**PDB header:**cytokine **Chain:** A: **PDB Molecule:**stress response peptide-2; **PDBTitle:** cytokine-like stress response peptide-2 in manduca sexta
13	c2w50B_	Alignment		6.9	24	**PDB header:**hormone **Chain:** B: **PDB Molecule:**armet-like protein 1; **PDBTitle:** n-terminal domain of human conserved dopamine neurotrophic factor2 (cdnf)

14	c2b3yB_	Alignment		5.9	32	**PDB header:**lyase **Chain:** B: **PDB Molecule:**iron-responsive element binding protein 1; **PDBTitle:** structure of a monoclinic crystal form of human cytosolic aconitase2 (irp1)